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ATTORNEY DOCKET NO.

09/057,675

FILING DATE

FIRST NAMED INVENTOR

APPLICATION NO.

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07/20/99

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NON-FINAL OA

CLIFFORD KENT WEBER FER THOMAS JEFFERSON UNIVERSITY OFFICE OF UNIVERSITY COUNSEL 1020 WALNUT STREET SUITE 620 PHILADELPHIA PA 19107-5587

EXAMINER ...

LEE, G

ART UNIT

PAPER NUMBER

1632

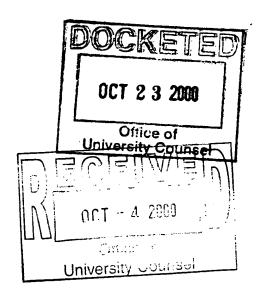
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DUE: 12/27/00

FINAL: 3/27/01

PI ase find below and/or attached an Office communication concerning this application or proceeding.

Commissioner of Patents and Trademarks



Office Action Summary

Application No. 09/357,675

Applicant(s)

Croce, Carlo M.

Examiner

Gai (Jennifer) Mi Lee

Group Art Unit 1632

Responsive to communication(s) filed on	
This action is FINAL.	·
☐ Since this application is in condition for allowance except for formal matters, in accordance with the practice under Ex parte Quay\835 C.D. 11; 453 O.G.	prosecution as to the merits is closed . 213.
A shortened statutory period for response to this action is set to expireonger, from the mailing date of this communication. Failure to respond within the application to become abandoned. (35 U.S.C. § 133). Extensions of time may be 37 CFR 1.136(a).	ne period for response will cause the
Disposition of Claim	
	is/are pending in the applicat
Of the above, claim(s) 4-9 and 12	is/are withdrawn from consideration
Claim(s)	is/are allowed.
X Claim(s) <u>1-3, 10, 11, and 13-16</u>	is/are rejected.
☐ Claim(s)	is/are objected to.
☐ Claims	are subject to restriction or election requirement.
Application Papers See the attached Notice of Draftsperson's Patent Drawing Review, PTO-9 The drawing(s) filed on is/are objected to by the The proposed drawing correction, filed on is	Examiner. approveddisapproved. § 119(a)-(d). uments have been ureau (PCT Rule 17.2(a)).
🔀 Acknowledgement is made of a claim for domestic priority under 35 U.S.C	C. § 119(e).
Attachment(s) Notice of References Cited, PTO-892 Information Disclosure Statement(s), PTO-1449, Paper No(s). Interview Summary, PTO-413 Notice of Draftsperson's Patent Drawing Review, PTO-948 Notice of Informal Patent Application, PTO-152	
SEE OFFICE ACTION ON THE FOLLOWIN	NG PAGES

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DETAILED ACTION

Applicant's election of Group I, Claims 1-3 and 10-16 without traverse in Paper No. 6 is acknowledged.

Please <u>note</u> that it appears that claim 12, directed to a pharmaceutical composition comprising a therapeutically effective amount of the Nit1 protein, had been inadvertently and improperly placed in the Invention of Group I rather than in the Invention of Group II. In light of the compact prosecution, claim 12 has been properly <u>rejoined</u> with the Invention of Group II (claims 4-6 and 13-14), drawn to a protein and uses thereof, which would properly encompass a pharmaceutical composition comprising a therapeutically effective amount of the Nit1 protein.

Claims 4-9 and 12 are withdrawn from further consideration by the examiner, 37 CFR 1.142(b), as being drawn to a non-elected Inventions. It is noted that in the <u>Response</u> section of the Restriction, Applicants indicate that claims 4-9 have been canceled. However, cancellation of a claim must take place <u>In the Claims</u> section of the Response to a restriction requirement. As such, claims 4-9 has not been canceled.

Claims 1-3, 10-11 and 13-16 are under examination only in so far as the claimed invention is drawn to the elected invention of a nucleic acid and methods of gene therapy using such.

Claims 1-3, 10-11 and 13-16 are currently under examination.

Priority

Applicant has not complied with one or more conditions for receiving the benefit of an earlier filing date under 35 U.S.C.119 (e) as follows:

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An application in which the benefits of an earlier application are desired must contain a specific reference to the prior application(s) in the first sentence of the specification (37 CFR 1.78).

Claim Objections

Claim 11 is objected to because of the following informalities: On line 2, a space should be inserted between Nit1 protein. Appropriate correction is required.

Claims 13- 15 are objected to because of the following informalities: The claims have not been amended to read on only the elected invention. Appropriate correction is required.

Claim Rejections - 35 USC § 112

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

Written Description

Claims 1-3, 10-11 and 13-16 are rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention. Applicant is referred to the interim guidelines on written description published December 21, 1999 in the Federal Register at Volume 64, Number 244, pp. 71440-71442 (also available at www.uspto.gov).

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"applicant must convey with reasonable clarity to those skilled in the art that, as of the filing date sought, he or she was in possession of the invention. The invention is, for purposes of the 'written description' inquiry, whatever is now claimed." Vas-Cath Inc. v. Mahurkar, 19USPQ2d at 1117. The specification does not "clearly allow persons of ordinary skill in the art to recognize that [he or she] invented what is claimed." Vas-Cath Inc. v. Mahurkar, 19USPQ2d at 1116.

The instant specification does not contain a written description of the invention in such full, clear, concise, and exact terms or in sufficient detail that one skilled in the art can reasonably conclude that applicant had possession of the claimed invention at the time of filing.

The claims are drawn to a purified NIT1 gene. In particular, wherein the gene is a human gene or mouse gene. The claims are also directed to an isolated nucleic acid of less than 100 kb comprising a nucleotide sequence encoding a Nit1 protein wherein the Nit1 protein is a human Nit1 protein (claims 10-11). The claims are further drawn to a method of treating or preventing a disease or disorder in a subject comprising administering to said subject a therapeutically effective amount of a molecule that inhibits or enhances Nit1 protein function (claims 13-14) or a method of gene therapy for treating or preventing a disease or disorder in a subject by using a vector containing the NIT1 gene coding sequence (claims 16). Claim 15 is drawn to a method of diagnosing or screening for the presence of or a predisposition for developing a disease or disorder in a subject comprising detecting one or more mutations in NIT1 DNA or RNA derived

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from the subject in which the presence of said one or more mutation indicates the presence of the disease or disorder or a predisposition for developing the disease or disorder.

The specification discloses an isolated cDNA sequence, SEQ ID NO: 1, which encodes a predictive polypeptide sequence. Absent evidence to the contrary, the *NIT1* gene elected for examination is deemed to be an incomplete cDNA. Because the cDNAs that correspond to the SEQ ID NO: 1 mentioned in the specification are not full-length, a sequence prepared from undefined parts of a cDNA clone will not comprise the entire coding region of any particular gene, nor is it clear the partial sequence is even in frame to encode a polypeptide. The claims, as written, however, encompass polynucleotides which vary substantially in length and also in nucleotide composition. The broadly claimed genus additionally, encompasses *NIT1* genes, as well as genes incorporating only portions of the disclosed sequence.

The instant disclosure of a single species of nucleic acid does not adequately describe the scope of the claimed genus, which encompasses a substantial variety of subgenera including full-length genes. A description of a genus of cDNAs may be achieved by means of a recitation of a representative number of cDNAs, defined by nucleotide sequence, falling within the scope of the genus or of a recitation of structural features common to members of the genus, which features constitute a substantial portion of the genus. *Regents of the University of California v. Eli Lilly & Co.*, 119 F3d 1559, 1569, 43 USPQ2d 1398, 1406 (Fed. Cir. 1997). The instant specification fails to provide sufficient descriptive information, such as definitive structural or functional features of the claimed genus of polynucleotides. There is no description of the conserved regions which

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are critical to the structure and function of the genus claimed. The specification proposes to discover other members of the genus by using a comparison with nitrilase and Fhit homologs which are encoded as fusion proteins in *D. melanogaster* and *C. elegans*. There is no description, however, of the sites at which variability may be tolerated and there is no information regarding the relation of structure to function. Structural features that could distinguish the compounds in the genus from others excluded are missing from the disclosure. Furthermore, the prior art does not provide compensatory structural or correlative teachings sufficient to enable one of skill to isolate and identify the polynucleotides encompassed and no identifying characteristic or property of the instant polynucleotides is provided such that one of skill would be able to predictably identify the encompassed molecules as being identical to those instantly claimed.

The specification further fails to identify and describe the 5' and 3' regulatory regions and untranslated regions essential to the function of the claimed invention, which are required since the claimed invention currently encompasses the gene. The art indicates that the structures of genes with naturally occurring regulatory elements and untranslated regions is empirically determined (Harris et al. J. of The Am Society of Nephrology 6:1125-33, 1995; Ahn et al. Nature Genetics 3(4):283-91, 1993; and Cawthon et al. Genomics 9(3):446-60, 1991). Therefore, the structure of these elements is not conventional in the art and skilled in the art would therefore not recognize from the disclosure that applicant was in possession of the genus of nucleic acid, including genes, comprising SEQ ID NO: 1.

Since the disclosure fails to describe the common attributes or characteristics that identify members of the genus, and because the genus is highly variant, the disclosure of specific nucleotide sequences and the ability to screen, is insufficient to describe the genus. One of skill in the art would reasonably conclude that the disclosure fails to provide a representative number of species to describe and enable the genus as broadly claimed.

Enablement

Claims 1-3 and 10-11 are rejected under 35 U.S.C. 112, first paragraph, because the specification, while being enabling for an isolated cDNA of a plant nitrilase NIT1, does not reasonably provide enablement for any and all nitrilase DNA (NIT1) of any species nor any isolated nucleic acid of less than 100 kb comprising a nucleotide sequence encoding a Nit1 protein. The specification does not enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make the invention commensurate in scope with these claims.

The claims are directed to any purified NIT1 gene wherein said gene is a human gene or mammalian gene (Claims 1-3). The claims are further directed to any isolated nucleic acid of less than 100 kb comprising a nucleotide sequence encoding a Nit1 protein wherein said protein is a human Nit1 protein (claims 10-11).

The specification discloses that human and murine NIT1 genes were cloned and characterized. Their exon-intron structure, their patterns of expression, and their alternative



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mRNA processing were determined and the tissue specificity of expression of murine FHIT and NIT1 genes were asserted to be nearly identical (page 2). The specification further discloses that the human and mouse NIT1 genes are members of an uncharacterized mammalian gene family with homology to bacterial and plant nitrilases, enzymes which cleave nitriles and organic amides to the corresponding carboxylic acids plus ammonia (page 3). Figure 1 teaches a sequence comparison of human, murine, D. melanogaster and C. elegans Nit1 and Fhit1 proteins. Figure 6 sets forth a highly conserved sequence of human, murine, D. melanogaster and C. elegans NIT1 gene (SEQ ID NO: 1). The specification further discloses that the mouse and human Nit1 amino acid sequences were 90% identical; the human Nit1 amino acid sequence was 58% similar and 50% identical to the C. elegans nitrilase domain and 63% similar and 53% identical to the D. melanogaster nitrilase domain (page 12 and Figure 1). However, the specification fails to teach or provide parameters, mechanistic characteristics, or classes of nitrilase domains for which one of skill in the art could reasonably predict that the NIT1 gene encodes a functional Nit1 protein which exists in any species and which can be utilized to practice the claimed invention without undue experimentation due to the unpredictability of the function of various plant and bacterial nitrilases known in the art, as well as the absence of guidance provided by the specification as to any NIT1 gene with functions in the same manner as the fusion gene, Fhit, of D. melanogaster and C. elegans.

Accordingly, in view of the quantity of experimentation necessary to determine the NIT1 gene of any species or an isolated nucleic acid encoding any Nit1 protein, the lack of direction or

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guidance provided by the specification as well as the absence of working examples with regards to the breadth of the claims directed to any NIT1 gene or any isolated nucleic acid that is less than 100 kb comprising a nucleotide sequence encoding a Nit1 protein, it would have required undue experimentation for one skilled in the art to make the claimed invention as broadly claimed.

Claims 13-16 are rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to enable one skilled in the art to which it pertains, or with which it is most nearly connected, to make and/or use the invention. The factors to be considered have been summarized as the quantity of experimentation necessary, the amount of direction or guidance presented, the presence or absence of working examples, the nature of the invention, the state of the prior art, the relative skill of those in the art, the predictability or unpredictability of the art and the breadth of the claims. *Ex Parte Forman*, (230 USPQ 546 (Bd Pat. App. & Int. 1986)).

The claims are drawn to a method of treating or preventing a disease or disorder in a subject comprising administering to said subject a therapeutically effective amount of a molecule that inhibits or enhances Nit1 function (claims 13 and 14) interpreted as directed to gene therapy (See page 1). In further embodiment, the claims are drawn to a method of diagnosing or screening for the presence of or a predisposition for developing a disease or disorder in a subject comprising detecting one or more mutations in NIT1 DNA. RNA or Nit1 protein derived from

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the subject (claim 15). The claims are also drawn to a method of treating or preventing a disease or disorder in a subject by using a vector containing the *NIT1* gene coding sequence (claim 16).

The claims are not enabled as the specification does not provide guidance as to the dosage amounts, dosage frequencies, modes of delivery, vectors for delivery, appropriate expression levels and targeting to supply any type of therapeutic treatment. The specification discloses that a tumor suppressor gene *FHIT* encompasses the common human chromosomal fragile site at 3p14.2 and numerous cancer cell bi-allelic deletions (page 2). The specification further discloses that in human and mouse, the nitrilase homologs and Fhit are encoded by two different genes, *FHIT* and *NIT1*, localized on chromosomes 3 and 1 in human, and 14 and 1 in mouse, respectively (page 2). The specification discloses that neither the *in vivo* function of Fhit nor the mechanism of its tumor suppressor activity is known but that analysis suggest that the enzyme-substrate complex is the active form that signals for tumor suppression (page 3). The specification further supports the unpredictability of Fhit function by stating that although the frequent loss of Fhit expression in several common human cancers is well documented, and results supporting its tumor suppressor activity have been reported, the role of Fhit in normal and tumor cell biology and its mechanism of its action *in vivo* is unknown (page 14).

While the specification teaches the skilled artisan how to determine the enzymatic function of the Nit1 protein in the claimed compositions only on the basis of similar homology to the Fhit fusion gene of *D. melanogaster* and *C. elegans*, it fails to provide guidance to the skilled artisan on how to use the claimed methods for <u>any</u> treatment of <u>any</u> disease or disorder. In

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particular, no protocol is described in the specification comprising administration of the polynucleotide or any other nucleic acid molecules that inhibit or enhance Nit1 function. The specification does not provide sufficient guidance as to the appropriate route of administration of any vectors for treating any diseases/disorders, the appropriate concentration of vectors, dosage, promoter to regulate expression for any treatment such that one of skill in the art could reproducibly, consistently, and effectively treat the patient in need thereof without undue experimentation.

Although the concept of gene therapy has potential, the realities of the parameters which will result in therapeutic benefit have not been achieved and are considered unpredictable. With regard to *in vivo* gene transfer, the specification provides no example or therapeutic methodology that would be encompassed within claims 13, 14 and 16. For example, Eck & Wilson (The Pharmacological Basis of Therapeutics, 1996) teach numerous factors complicate the gene delivery art which would not have been shown to be overcome by routine experimentation.

These include, the fate of the DNA vector itself (volume of distribution, rate of clearance into the tissues, etc.), the *in vivo* consequences of altered gene expression and protein function, the fraction of vector taken up by the target cell population, the trafficking of the genetic material within cellular organelles, the rate of degradation of the DNA, the level of mRNA produced, the stability of the mRNA produced, the amount and stability of the protein produced, and the protein's compartmentalization within the cell, or its secretory fate, once produced. These factors differ dramatically based on the vector used and the protein being produced, which cells

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are target cells, and the disease and/or host being treated. It is further noted that Eck and Wilson support the importance of tailoring a gene therapy vector and method to specific diseases and/or disorders. See page 82, column 1, first paragraph. Furthermore, Eck & Wilson et al. review the state of the art for gene therapy for inherited disorders and discloses that "[t]he level of protein function necessary to achieve complementation of the defect varies widely among genetic diseases." See page 78, column 2, 2nd paragraph.

In addition, while progress has been made in recent years for gene transfer *in vivo*, vector targeting to desired tissues *in vivo* continues to be unpredictable and inefficient as supported by numerous teachings available in the art. For example, Miller (1995, FASEB J., Vol. 9, pages 190-199) review the types of vectors available for *in vivo* gene therapy, and conclude that "for the long-term success as well as the widespread applicability of human gene therapy, there will have to be advances...targeting strategies outlined in this review, which are currently only at the experimental level, will have to be translated into components of safe and highly efficient delivery systems" (page 198, column 1). Deonarain (1998, Expert Opin. Ther. Pat., Vol. 8, pages 53-69) indicate that one of the biggest problems hampering successful gene therapy is the "ability to target a gene to a significant population of cells and express it at adequate levels for a long enough period of time" (page 53, first paragraph). Deonarain reviews new techniques under experimentation in the art which show promise but states that such techniques are even less efficient than viral gene delivery (see page 65, first paragraph under Conclusion section). Verma (Sept. 1997, Nature, Vol. 389, pages 239-242) reviews vectors known in the art for use in gene

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therapy and discusses problems associated with each type of vector. The teachings of Verma indicate a resolution to vector targeting has not been achieved in the art (see entire article). Verma also teaches appropriate regulatory elements may improve expression, but it is unpredictable what tissues such regulatory elements target (page 240, sentence bridging columns 2 and 3). Crystal (1995, Science, Vol. 270, page 404-410) also reviews various vectors known in the art and indicates that "among the design hurdles for all vectors are the need to increase the efficiency of gene transfer, to increase target specificity and to enable the transferred gene to be regulated" (page 409).

Specifically, the specification, on page 7, teaches only *in vitro* multiple tissue northern blots of *NIT1* cDNA probes. However, no further results are reported on the effectiveness of Nit1 protein function having any implication toward the treatment of any particular diseases or disorders. It is noted that, Orkin stress the importance of using relevant animal models for determining the effectiveness of therapeutic methodologies (p. 10 and 13). As such, the specification fails to provide any evidence which would provide a reasonable nexus to that of any particular diseases or disorders.

With regard to claim 15, directed to a method of diagnosing or screening for the presence of or a predisposition for developing a disease or disorder in a subject comprising detecting one or more mutations in NIT1 DNA or RNA derived from the subject in which the presence of said one or more mutations indicates the presence of the disease or disorder or a predisposition for developing the disease or disorder, the specification fails to teach or suggest any methodology or

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procedure for a method of diagnosing or screening of any mutations in NIT1 DNA or RNA correlating to any disease or disorder in any subject as embraced by the claim. The specification only discloses that the pattern of Nit1 expression was almost identical to the pattern of the expression of Fhit (Fig. 2A), supporting the hypothesis that the proteins may act in concert or participate in the same pathway (page 14, lines 1-4). However, the specification fails to discuss any methods of screening or diagnosing of any disease or disorder in any subject comprising detecting any mutations in NIT1 DNA or RNA in which the presence of any said mutations would indicate the presence of any disease or disorder. The specification further fails to indicate that any mutations in NIT1 DNA or RNA would even correlate to any disease or disorder. Thus, it would be unpredictable for one of skill in the art to identify mutations of NIT1 DNA or RNA which would result in any disease or disorder as embraced by the claimed invention.

If claims 13-16 should be overcome by applicants arguments and/or evidence, claim 13-16 would be limited to use and specific therapeutic function of the exemplified Nit 1 nucleic acid sequence.

Accordingly, in view of the unpredictable and undeveloped state of the art, the lack of guidance or working examples which demonstrate or correlate to any therapeutic effect of the claimed methods, including the identification of any Nit1 mutations, and the breadth of the claims, the specification fails to teach any nucleic acid sequences which "enhance or inhibit" a Nit 1 protein as embraced by the claims directed to a method of treating or preventing any disease or disorder in any subject, including any are disease relevant Nit 1 mutations.

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The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claims 1-3, 10-11 and 13-15 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

Claims 1, 10 and 11 are vague and indefinite for its recitation of "NIT1 gene" because it is unclear from the specification as to what is encompassed within the claims as to "NIT1" gene since Figure 6 (SEQ ID NO: 1) is "NITD". Is NIT1 the same as NITD? The metes and bounds of the claim cannot be determined. Note that claims 2-3 depends from claim 1.

Claims 13 and 14 are incomplete. While all of the technical details of method need not be recited, the claims should include enough information to clearly and accurately describe the invention and how it is practiced. The method of claims 13 and 14 are missing process steps.

The method step needs to correlate to the preamble because it is unclear as to how inhibiting Nit1 function or enhancing Nit1 function would treat or prevent any disease or disorder.

Claim 15 is incomplete. While all of the technical details of method need not be recited, the claims should include enough information to clearly and accurately describe the invention and how it is practiced. The method of claim 15 is missing process steps. The method step needs to correlate to the preamble because it is unclear as to how mere detecting one or more mutations in NIT1 DNA or RNA would diagnose or screen for the presence of or a predisposition for developing any disease or disorder.

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Claim 16 is incomplete. While all of the technical details of method need not be recited, the claims should include enough information to clearly and accurately describe the invention and how it is practiced. The method of claim 16 is missing process steps. In addition, the method step needs to correlate to the preamble because it is unclear as to how using a vector containing *NIT1* gene coding sequence would treat or prevent any disease or disorder.

Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless --

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

Claims 1 and 10 are rejected under 35 U.S.C. 102(b) as being anticipated by Hillebrand et al (May 8, 1996) Gene, Vol. 170 (2): 197-200.

The claims are drawn to <u>any and all</u> purified NIT1 gene (claim 1) or <u>any isolated nucleic</u> acid of less than 100 kb comprising a nucleotide sequence encoding <u>any Nit1 protein</u> (claim 10).

Hillebrand et al disclose that a <u>full-length genomic clone</u> encoding the **complete** cluster of the *At* nitrilases 1-3 (NIT 1-3), including the respective promoter regions, has been isolated and sequenced. Thus, Hillebrand et al clearly anticipate claims 1 and 10 of the instant invention.

Conclusion

Claims 2-3, 11 and 13-16 appear to be free of the cited prior art of record because the cited prior art of record fails to teach or suggest a purified human or mammalian NIT1 gene as

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well as a method of treating, preventing, diagnosing or screening any disease or disorder using the same. However, these claims are subject to other rejections.

No claims are allowed.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Gai (Jennifer) Mi Lee, whose telephone number is 703-306-5881. The examiner can normally be reached on Monday-Thursday from 8:30 to 5:00 (EST). The examiner can also be reached on alternate Fridays.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Karen Hauda, can be reached on 703-305-6608. The FAX phone numbers for group 1600 are 703-308-4242 and 703-305-3014.

An inquiry of a general nature or relating to the status of the application should be directed to the group receptionist whose telephone number is 703-308-0196.

Gai (Jennifer) Lee Patent Examiner Art Unit 1600

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	Notice of References Cited 09/357,675 Croce, Carlo M.						
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Application No.

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	Hillebrand et al., Gene, Vol.	170(2): 197-20	00.				5/1996
	•	170(2): 197-20	00.				5/1996

U.S. DEPARTMENT OF COMMERCE - Patent and Trademark Office Application No.

NOTICE OF DRAFTSPERSON'S PATENT DRAWING REVIEW

1. DRAWINGS: 37 CFR 1.84(1) Blockink, Color (1) 17 (1) 37 (1) 18 (b) Fig(s) Percil and non-block-lick, one permitted, Fig(s) Percil and non-block-lick, one permitted, Fig(s) Photographic double-weight paper), Fig(s) 1. full-tone set is required. Fig(s) Poor quality (half-tone), Fig(s) 3. TYPE OF PAPER, 37 CFR 1.84(e) Paper on disable, strong, white, and darabla, Fig(s) Fig(s) 1. region of the block one permitted fig(s) Paper on disable, strong, white, and darabla, Fig(s) Fig(s) 1. region of the block one permitted fig(s) Paper on disable, strong, white, and darabla, Fig(s) Fig(s) 1. CHARACTER OF LINES, NUMBERS, & LETTERS. 37 CFR 1.84(s) Fig(s) 1. SHADING, 37 CFR 1.84(c) Lices, numbers & letters not uniformly thick and well defined; clean, durable, and black (poor line quality). Fig(s) Solid black areas pale. Fig(s) Numbers and reference characters not oriented in the same difference characters may be at least 37 CFR 1.84(s) Fig(s) Top (7) Left (Right (R) Fig(s) Fig(s) Solid blac	The drawing(s) filed (insert date) 7-70-97 are: A. approved by the Draftsperson under 37 CFR 1.84 or 1.152. B. objected to by the Draftsperson under 37 CFR 1.84 or 1.152 for the results of the content of the conten	easons indicated below. The Examiner will require ust be sumitted according to the instructions on the back of this notice
DRAWINGS: 37 CFR 1.84(b) Acceptable categories of drawings		
Marries not acceptable Fig(s) Top (T): Right (R) Bottom (B): 6. VIEWS. 37 CFR 1.84(h). REMINDER: Specification may require revision to correspond to drawing changes: Partial views. 37 CFR 1.84(h)(2): Brackets needed to show figure as one entity. Fig(s) Enlarged view not labeled separately or properly. Fig(s) Enlarged view not labeled separately or properly. Fig(s) Toc (I/8 igch) in height. 37 CFR 1.84(q) Lead lines cross each other. Fig(s) Sheets not numbered consecutively, and in Arabic numerals beginning with number 1. Sheet(s) 15. NUMBERING OF VIEWS. 37 CFR 1.84(u) Views not numbered consecutively, and in Arabic numerals, beginning with number 1. Fig(s) Enlarged view not labeled separately or properly. Fig(s) To SECTIONAL VIEWS. 37 CFR 1.84 (h)(3) Hatching not indicated for sectional portions of an object. Fig(s) Sectional designation should be noted with Arabic or Roman numbers. Fig(s) To SIGN DRAWINGS. 37 CFR 1.152 Surface shading not used for color contrast. Fig(s) Solid black shading not used for color contrast.	1. DRAWINGS. 37 CFR 1.84(a)? Acceptable categories of drawings: Black,ink., Colordy 2019/810 Mon 201 Color drawings are not acceptable until petition is granted. Fig(s) Pencil and non.black.tok.not/permitted. Fig(s) Pencil and non.black.tok.not/permitted. Fig(s) Pencil and non.black.tok.not/permitted. Fig(s) Photographs. 37 CFR 1.84 (b) I full-tone set is required. Fig(s) Photographic double-weight paper). Fig(s) Foor quality (half-tone). Fig(s) 3. TYPE OF PAPER. 37 CFR 1.84(g) Paper-not flexible, strong, white, and durables. Fig(s) Erasures; alterations; overwritings, interdineations; Mylar, velum paper is not acceptable (too thin). Fig(s) 4. SIZE OF PAPER. 37 CFR 1.84(f): Acceptable sizes: 21.0 cm by 29.7 cm (DIN size A4) 21.6 cm by 27.9 cm (8 1/2 x 11 inches) Sheet(s) Drawings sheets not an acceptable size. Fig(s) 5. MARGINS. 37 CFR 1.84(g): Acceptable margins: 882-() [7] Drawings sheets not an acceptable size. Fig(s) Size: A4 Size: A4 Size. Bottom 1.0 cm SIZE: A4 Size. Bottom 1.0 cm SIZE: A4 Size.	8.5 ARRANGEMENT OF. VIEWS; 37 CFR 1.84(i) Words do not appear on a horizontal, left-to-right fashion when page is either upright or turned so that the top becomes the right side, except for graphs. Fig(s) 9. SCALE. 37 CFR 1.84(k): Scale not large enough to show mechanism without crowding when drawing is reduced in size to two-thirds in reproduction. Fig(s) 10. CHARACTER OF LINES, NUMBERS, & LETTERS. 37 CFR 1.84(i) Lines, numbers & letters not uniformly thick and well defined; clean, durable, and black (poor line quality). Fig(s): Solid black areas pale. Fig(s): Solid black areas pale. Fig(s): Shade lines, pale; rough and blurred. Fig(s): 12. NUMBERS, LETTERS, & REFERENCE CHARACTERS. 37 CFR 1.84(p): Numbers and reference characters not plain and legible. Fig(s): Figure legends are poor. Fig(s): Numbers and reference characters not oriented in the same direction as the view. 37 CFR 1.84(p)(1). English alphabet not used. 37 CFR 1.84(p)(2)
COMMENTS	Margins not acceptable: Fig(a) Top (T): Right (R) Bottom (B): 6. VIEWS. 37 CFR 1.84(h). REMINDER: Specification may require revision to correspond to drawing changes. Partial views. 37 CFR 1.84(h)(2): Brackets needed to show figure as one entity. Fig(a) Views not labeled separately or properly. Fig(a) Enlarged view not labeled separately or properly. Fig(a) SECTIONAL VIEWS. 37 CFR 1.84 (h)(3) Hatching not indicated for sectional portions of an object. Fig(s) Sectional designation should be noted with Arabic or Roman numbers. Fig(a)	Numbers, letters and reference characters must be at least 3.2 cm (1/8 ligch) in height. 37 CFR 1.84(p)(3) 13. LEAD LINES. 37 CFR 1.84(q) Lead lines cross each other. Fig(s) Lead lines missing. Fig(s) 14. NUMBERING OF SHEETS OF DRAWINGS. 37 CFR 1.84(t) Sheets not numbered consecutively, and in Arabic numerals beginning with number 1. Sheet(s) 15. NUMBERING OF VIEWS. 37 CFR 1.84(u) Views not numbered consecutively, and in Arabic numerals, beginning with number 1. Fig(s) 16. CORRECTIONS. 37 CFR 1.84(w) Corrections not made from prior PTO-948 dated 17. DESIGN DRAWINGS. 37 CFR 1.152 Surface shading shown not appropriate. Fig(s) Solid black shading not used for color contrast.

DATE 8-13.99 TELEPHONE NO.

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PTO/SB/21 (08-00)

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TRANSMITTAL FORM

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Total Number of Pages in This Submission

Application Number	09/357,675
Filing Date	July 20, 2001
First Named Inventor	CROCE, Carlo
Group Art Unit	1632
Examiner Name	LEE, G.
Attorney Docket Number	CRO01-NP001

	·	ENCLOSURES (chec	k all that apply)
X Fee Transmittal Fo	rm	Assignment Papers (for an Application)	After Allowance Communication to Group
Fee Attache	ed	Drawing(s)	Appeal Communication to Board of Appeals and Interferences
X Amendment / Repl	у	Licensing-related Papers	Appeal Communication to Group (Appeal Notice, Brief, Reply Brief)
After Final		Petition	Proprietary Information
Affidavits/declaration(s)		Petition to Convert to a Provisional Application	Status Letter
X Extension of Time	Request	Power of Attorney, Revocation Change of Correspondence Address	Other Enclosure(s) (please identify below):
Express Abandonn	nent Request	Terminal Disclaimer Request for Refund	
Information Disclos	sure Statement	CD, Number of CD(s)	_
Certified Copy of P Document(s)	riority	Remarks	***************************************
Response to Missii			ich ,
Response to	Missing Parts FR 1.52 or 1.53		PECH CENTER SOLUTION
	SIGNATU	URE OF APPLICANT, ATTORNEY, OF	RAGENT & 2
Firm or Individual name	Clifford K. Wel	ber) FEC 12900
Signature	Cliffind X. W	Velver	
Date	March 27, 2001	1	

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FEE TRANSMITTAL for FY 2001

Patent fees are subject to annual revision.

TOTAL AMOUNT OF PAYMENT

(\$) 455.00

Co	omplet if Known	70	M
Application Number	09/357,675	田山	
Filing Date	July 20, 1999	20	
First Named Inventor	CROCE, Carlo	9 9	
Examiner Name		129	r.
Group Art Unit		10	
Attorney Docket No.	CRO01-NP001		

METHOD OF PAYMENT FEE CALCULATION (continued)				
1. The Commissioner is hereby authorized to charge indicated fees and credit any overpayments to:	3. ADDITIONAL FEES			
indicated fees and credit any overpayments to: Deposit	Large Small			
Account Number 50-0491	Entity Entity Fee Fee Fee Fee Fee Description	Fee Paid		
Denosit	Code (\$) Code (\$)	1001010		
Account Name Thomas Jefferson University	105 130 205 65 Surcharge - late filing fee or oath			
Charge Any Additional Fee Required Under 37 CFR 1.16 and 1.17	127 50 227 25 Surcharge - late provisional filing fee or cover sheet			
Applicant claims small entity status.	139 130 139 130 Non-English specification			
See 37 CFR 1.27	147 2,520 147 2,520 For filing a request for ex parte reexamination			
2. Payment Enclosed: Check Credit card Money Other	112 920* 112 920* Requesting publication of SIR prior to Examiner action			
FEE CALCULATION	113 1,840* 113 1,840* Requesting publication of SIR after Examiner action			
	115 110 215 55 Extension for reply within first month			
1. BASIC FILING FEE Large Entity Small Entity	116 390 216 195 Extension for reply within second month			
Fee Fee Fee Fee Description	117 890 217 445 Extension for reply within third month	\$455.00		
Code (\$) Code (\$) Fee Paid 101 710 201 355 Utility filing fee	118 1,390 218 695 Extension for reply within fourth month			
106 320 206 160 Design filling fee	128 1,890 228 945 Extension for reply within fifth month			
107 490 207 245 Plant filing fee	119 310 219 155 Notice of Appeal			
108 710 208 355 Reissue filing fee	120 310 220 155 Filing a brief in support of an appeal			
114 150 214 75 Provisional filing fee	121 270 221 135 Request for oral hearing			
	138 1,510 138 1,510 Petition to institute a public use proceeding			
SUBTOTAL (1) (\$) 0.00	140 110 240 55 Petition to revive - unavoidable			
2. EXTRA CLAIM FEES	141 1,240 241 620 Petition to revive - unintentional			
Extra Claims below Fee Paid	142 1,240 242 620 Utility issue fee (or reissue)			
Total Claims20** = X \$9.00 = \$0.00	143 440 243 220 Design issue fee			
Independent - 3** = X \$40.00 = \$0.00	144 600 244 300 Plant issue fee			
Multiple Dependent	122 130 122 130 Petitions to the Commissioner			
	123 50 123 50 Processing fee under 37 CFR 1.17(q)			
Large Entity Small Entity Fee Fee Fee Fee Description	126 180 126 180 Submission of Information Disclosure Stmt			
Code (\$) Code (\$) 103 18 203 9 Claims in excess of 20	581 40 581 40 Recording each patent assignment per property (times number of properties)			
102 80 202 40 Independent claims in excess of 3	146 710 246 355 Filing a submission after final rejection (37 CFR § 1.129(a))			
104 270 204 135 Multiple dependent claim, if not paid 109 80 209 40 ** Reissue independent claims	149 710 249 355 For each additional invention to be examined (37 CFR § 1.129(b))			
over original patent 110 18 210 9 ** Reissue claims in excess of 20	179 710 279 355 Request for Continued Examination (RCE)			
and over original patent	169 900 169 900 Request for expedited examination of a design application			
SUBTOTAL (2) (\$)0.00	Other fee (specify)			
**or number previously paid, if greater; For Reissues, see above	*Reduced by Basic Filing Fee Paid SUBTOTAL (3) (\$) 455	5.00		

SUBMITTED BY					Complete (if applicable)	
Name (PrintlType)	Clifford K. Weber	Registration No. (Attorney/Agent)	42,215	Telephone	(215) 503-0757	
Signature	Cliffed F. Webe.			Date	March 27, 2001	